

Fig. 1

1 TCAATCAACGAGGCGTCCGTGCGACACAGGAGGAAATCCAATGAGCGGAAAACCTGGCTTA 60  
M S G K L A Y

61 CGTTACAGGCGGGATGGGCGGTATCGGCACCTCAATTTGCCAGCGCCTGGCCAAAGATGG 120  
V T G G M G G I G T S I C Q R L A K D G

121 CTTTCGCGTGGTGGCAGGCTGCGGCCCCAGCCGCAATTACCAGCAATGGCTGGATGAACA 180  
F R V V A G C G P S R N Y Q Q W L D E Q

181 GGCGGCGCAGGGCTATACGTTCTACGCGTCAGTGGGCAACGTGTCCGATTGGGAGTCCAC 240  
A A Q G Y T F Y A S V G N V S D W E S T

241 GGTAGAAGCATTGAGCGCGTCAAGCGGGACATGGGCCCGGTGATGTGCTGGTCAACAA 300  
V E A F E R V K R D M G P V D V L V N N

301 CGCGGGCATCACCCGCGACGGCCTGTTCCGCAAGATGAGCGCCGACGACTGGCGCGCGGT 360  
A G I T R D G L F R K M S A D D W R A V

361 CATCGACACCAACCTGAACAGCCTCTTCAACGTGACCAAAGCAGGTGATCGACGACATGGT 420  
I D T N L N S L F N V T K Q V I D D M V

421 CGAGCGCCAGTGGGGCCGCATCGTCAACATCAGCTCGGTGAACGGGCAGAAGGGGCGAGTT 480  
E R Q W G R I V N I S S V N G Q K G Q F

481 CGGCCAGACGAACTATTCCACGGCGAAGGCGGGCATCCATGGCTTCACCATGGCGCTGGC 540  
G Q T N Y S T A K A G I H G F T M A L A

541 GCAGGAAGTGGCCAGCAAGGGCATCACGGTCAACACGGTGTGCGCGGGCTACATCGGCAC 600  
Q E V A S K G I T V N T V S P G Y I G T

601 GGACATGGTTCGCGCCATCCGTCCGGACGTGCTGGAAAAGATCGTCGCCACCATTCCGGT 660  
D M V R A I R P D V L E K I V A T I P V

661 GCGCCGCTGGGCACGCCGGAGGAAATCGCGTCCATCACGTGCTGGCTGGCCTCGGATGA 720  
R R L G T P E E I A S I T S W L A S D E

721 GTCTGGGTTTTGACGGGCGCGGACTTCTCGCTCAACGGCGGCCTGCATATGGGCTGAAG 780  
S G F S T G A D F S L N G G L H M G \*

781 CATCGCGGGCCGCCACGAGCGGCCCCGCCGCGCGGGCGGCCTCGGGGAGAGGGCCGTCC 840

841 GGCATTACACTTACCCTTATCCGAAGTCTTAGAGATCGCCCGATCCGGGGACAACCATGA 900

Fig. 2

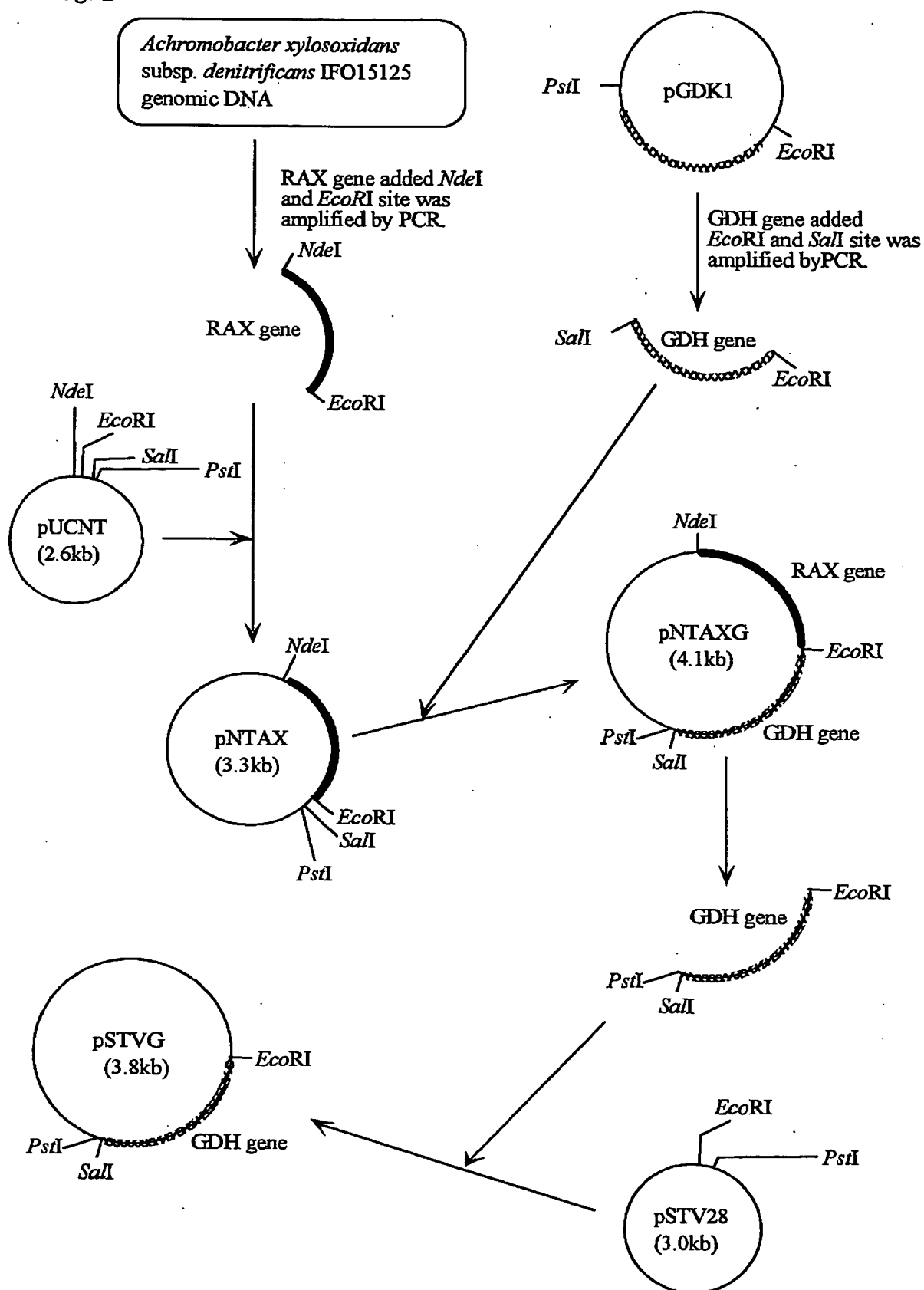


Fig. 3

